Combining Covariate Adjustment with Group Sequential Designs to Improve Randomized Trial Efficiency



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Joint work with

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Covariate Adjustment

Covariate adjustment is a statistical analysis method w	ith
high potential to improve precision for many trials.	

- **Pre-planned** adjustment for baseline variables when estimating **average treatment effect**.
- Estimand is same as when using unadjusted estimator (e.g., difference in means).
- Goal: avoid making any model assumptions beyond what's assumed for unadjusted estimator (robustness to model misspecification).

(e.g., Koch et al., 1998; Yang and Tsiatis, 2001; Rubin and van der Laan, 2008; Tsiatis et al., 2008; Moore and van der Laan, 2009b,a; Zhang, 2015; Jiang et al., 2018; Benkeser et al., 2020)

- A commonly used type of clinical trial design that involves **pre-planned interim analyses**
 - where the trial can be **stopped early** for efficacy or futility.
- Prevalent in confirmatory clinical trials for ethical and efficiency reasons as they potentially save time and resources by allowing early termination of the trial.

Problem Setting

- Combination of covariate adjustment and group sequential designs has the potential to offer the benefits of both methods:
 - using covariate adjusted estimators at interim and final analyses of a group sequential design.
- Several challenges involved in combining these two approaches.

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- 2 The uncertainty at the design stage about the amount of precision gain and corresponding sample size reduction.
 - Proposals have been made to use external trial data to estimate the precision gain.
 - Nevertheless, an incorrect projection of a covariate's prognostic value risks an over- or underpowered future trial.
 - ☐ This is also an obstacle in trials without interim analyses.

Endpoints, Estimands and Estimators

- The proposal works for all types of common outcomes
 - e.g., continuous, binary, ordinal, and time-to-event
- The proposal accommodates **any estimand**, including
 - risk difference, relative risk and odds ratio for binary outcomes,
 - difference in restricted mean survival times and relative risk for time-to-event outcomes.
 - ...
- The proposal will be applicable to **any (adjusted and unadjusted) estimator** as long as it is regular and asymptotically linear (RAL) and consistent for the estimand of interest.
 - e.g., G-computation estimator (as suggested in the recent FDA draft guidance on covariate adjustment)

Endpoints, Estimands and Estimators: Example

■ Primary endpoint: binary.

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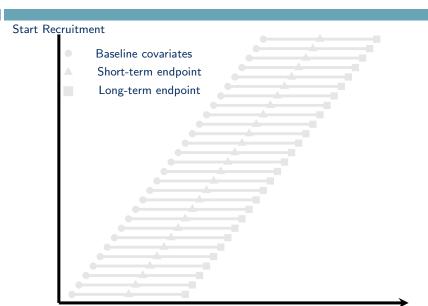
- Primary endpoint: binary.
- Estimand: $\theta = E(Y|A=1) E(Y|A=0)$.

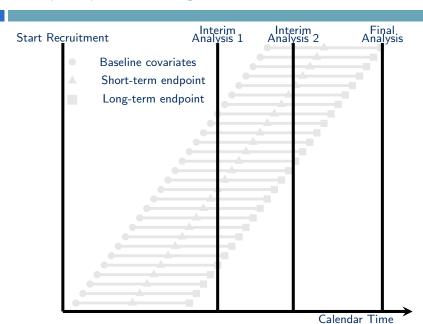
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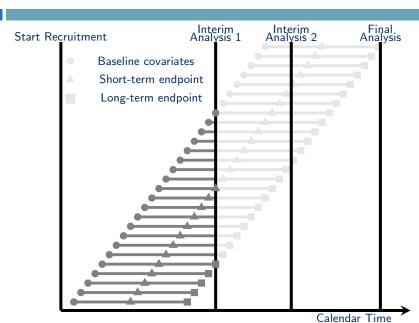
- Primary endpoint: binary.
- Estimand: $\theta = E(Y|A=1) E(Y|A=0)$.
- Estimator: G-computation/Standardization
 - 1 Fit logistic regression model for

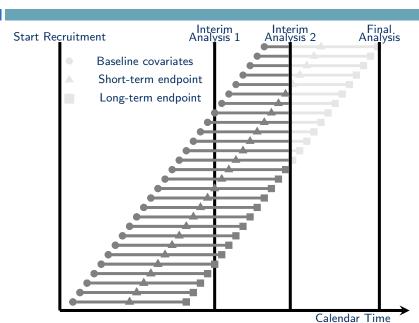
$$P(Y = 1|A, B) = logit^{-1}(\gamma_0 + \gamma_1 A + \gamma_2 B).$$

- Compute standardized estimators for treatment specific means
 - $\hat{E}(Y|A=1) = \frac{1}{n} \sum_{i=1}^{n} logit^{-1} (\hat{\gamma}_0 + \hat{\gamma}_1 + \hat{\gamma}_2 B_i)$
 - $\hat{E}(Y|A=0) = \frac{1}{n} \sum_{i=1}^{n} logit^{-1} (\hat{\gamma}_0 + \hat{\gamma}_2 B_i)$
- 3 Calculate $\hat{\theta} = \hat{E}(Y|A=1) \hat{E}(Y|A=0)$

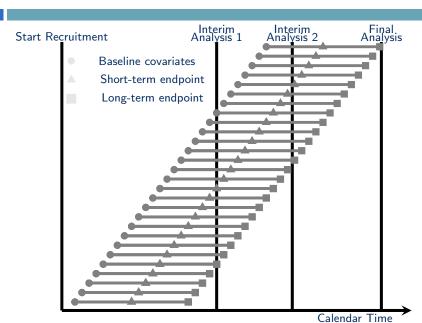








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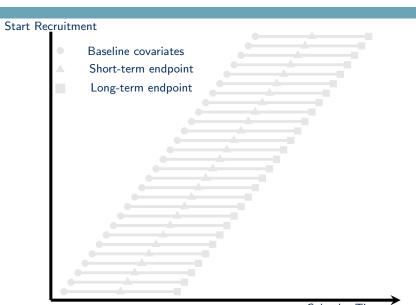
■ Entail analyzing the data at K different times t_1, \ldots, t_K .

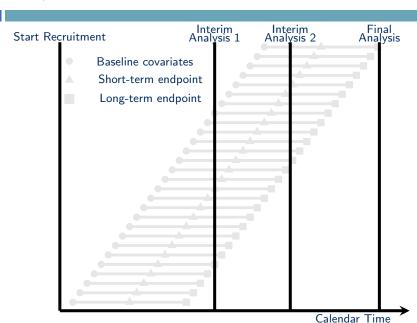
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- At each analysis time t_k :
 - \square Calculate an estimate $\hat{\theta}_{t_k}$.
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 - \square Compare the Z_k to some critical value for that analysis.
 - Allow to stop early for efficacy and/or futility.

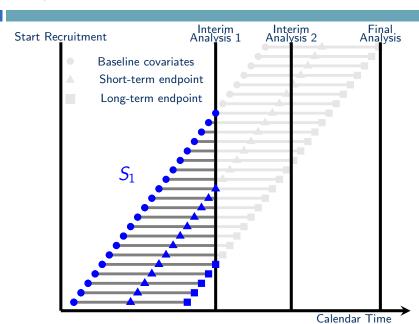
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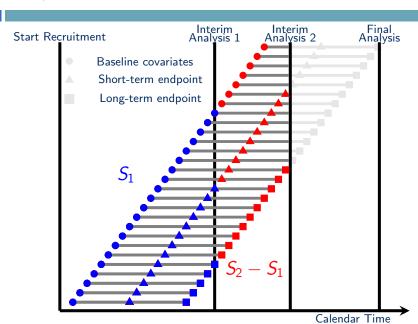
- A group sequential design can reduce the length of a Phase 3 trial
 - Reaching a conclusion sooner (Stopping for efficacy/futility)
 - ☐ Faster access to effective treatments
 - ☐ Faster dropping of ineffective/harmful treatment
 - Saving resources

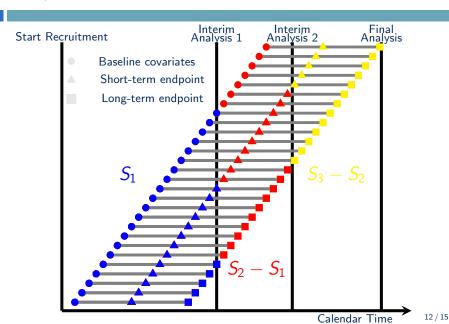
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☐ Saving resources
Multiple looks at accumulating data increase type I error
Lower significant thresholds needs to be used for the interim analyses.
There are a range of methods for defining the critical values for interim analyses.
(Pocock, 1977; O'Brien and Fleming, 1979; Lan and DeMets, 1983)











Group Sequential Designs: Incompatibility

- Unfortunately, a sequence of RAL estimators $(\hat{\theta}_{t_1}, \dots, \hat{\theta}_{t_K})$ do not necessarily have the independent increments property.
- This was for example shown for:
 - Estimators based on generalized estimating equations (Shoben and Emerson, 2014)
 - G-computation and TMLE estimators when working models are misspecified (Rosenblum et al., 2015)
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 - A long list of further counterexamples is provided by Jennison and Turnbull (1997) and Kim and Tsiatis (2020)
- Proposal: modifying any RAL estimator so that it has the independent increments property and also has equal or smaller variance than the original estimator.

Proposal: Motivation

- **Goal:** Obtain at each analysis time t_k an estimator θ_{t_k} that
 - 1 is consistent for θ ,
 - 2 is asymptotically linear,
 - is asymptotically normal,
 - is asymptotically as or more precise as the original estimator $\widehat{\theta}_{t_k}$, and
 - 5 has the independent increments property.

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 - 4 is asymptotically as or more precise as the original estimator $\widehat{\theta}_{t_k}$, and
 - b has the independent increments property.
- We will focus on finding the **linear combination**

$$\widehat{\theta}_{t_k} - \sum_{k'=1}^{k-1} \lambda_{k'}^{(k)} (\widehat{\theta}_{t_k} - \widehat{\theta}_{t_{k'}})$$

with minimal variance.

Thank you for your attention!

Interested? https://doi.org/10.48550/arXiv.2201.12921

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